

253	16.4	0.4	0.4	21	1	ABX79874	EST polymorphic DN	326	15.8	0.4	20	1	ABD24432	AI652901-derived o
c 254	16.4	0.4	0.4	21	1	ACC79938	Thermus oshimai nu	327	15.8	0.4	20	1	ADF73043	Primer #1 of the i
255	16.4	0.4	0.4	21	1	ADN08366	373 cell transform	328	15.8	0.4	20	1	ADF73047	Primer #3 of the i
256	16.4	0.4	0.4	21	1	ADS87406	Human midkine (MDK	c 329	15.8	0.4	20	1	ADJ31745	Human amyloid beta
257	16.4	0.4	0.4	21	1	ADS87422	Human midkine (MDK	330	15.8	0.4	20	1	ADJ31781	Human amyloid beta
258	16.4	0.4	0.4	21	1	ADS87404	Human midkine (MDK	331	15.8	0.4	20	1	ADI38836	Human LIM domain k
259	16.4	0.4	0.4	51	1	ADC17063	Human single nucle	c 332	15.8	0.4	20	1	ADI38771	Human LIM domain k
260	16.2	0.4	0.4	20	1	ADO56495	Human cyclin-nucle	333	15.8	0.4	20	1	ADL70240	Murine p27Kip1 PCR
c 261	16.2	0.4	0.4	21	1	AAQ65930	Type II procollage	c 334	15.8	0.4	20	1	ADL61407	Human protein tyro
c 262	16.2	0.4	0.4	21	1	AAQ226192	Human polymorphic	335	15.8	0.4	20	1	ADOS7395	Kidney developmenp
c 263	16.2	0.4	0.4	21	1	AAZ40511	Human STE20-relate	336	15.8	0.4	20	1	ADN02369	PCR primer 2 used
c 264	16.2	0.4	0.4	21	1	AAA38134	Polynucleotide use	337	15.8	0.4	20	1	ADN03787	SFG probe for dete
c 265	16.2	0.4	0.4	21	1	ADJ72441	Human GPI20 antibo	c 338	15.8	0.4	21	1	AAT11643	WT1/EGR human TCC
c 266	16.2	0.4	0.4	21	1	ADJ13904	Human DNA probe us	c 339	15.8	0.4	21	1	AAV10466	Human osteosarcoma
c 267	16.2	0.4	0.4	21	1	ADS87436	Human midkine (MDK	340	15.8	0.4	21	1	AAA07691	Reverse primer for
c 268	16.2	0.4	0.4	30	1	ADN97224	AGC1 locus. Unide	c 341	15.8	0.4	21	1	AAZ44677	E. coli strain 015
c 269	16.2	0.4	0.4	31	1	RAI31038	Human single nucle	c 342	15.8	0.4	21	1	AAA10649	PCR primer #3 used
270	16	0.4	0.4	17	1	ABK02354	Human NOGO Amberzy	c 343	15.8	0.4	21	1	AAA59901	Human OP-1 Wt-1/Bg
271	16	0.4	0.4	17	1	ABL46891	Human GRID G-cleav	c 344	15.8	0.4	21	1	AAH62656	Synaptotagmin 5 po
272	16	0.4	0.4	17	1	ABL46750	Human GRID NCH rib	345	15.8	0.4	21	1	AAH62429	HERC1 polymorphism
273	16	0.4	0.4	17	1	ADC37824	Human AMLP1a scann	c 346	15.8	0.4	21	1	AAH88947	Human polymorphic
274	16	0.4	0.4	17	1	ADC37817	Human AMLP1a scann	347	15.8	0.4	21	1	ABK65628	Human single nucle
275	16	0.4	0.4	17	1	ADM54108	Human GRID mRNA su	348	15.8	0.4	21	1	ABK65740	Human single nucle
c 276	16	0.4	0.4	18	1	AAK67194	Human CD40 hairpin	c 349	15.8	0.4	21	1	ABK65740	Human polymorphism
277	16	0.4	0.4	18	1	AAF26668	Human Smad7 phosph	c 350	15.8	0.4	21	1	ABK99278	Hepatitis C virus
c 278	16	0.4	0.4	19	1	ADR75637	Human apolipoprote	c 351	15.8	0.4	21	1	ADD22525	Flatfish rhabdovir
c 279	16	0.4	0.4	19	1	ADR78255	Human apolipoprote	c 352	15.8	0.4	21	1	ADE78130	DNA oligo (SeqID 3
c 280	16	0.4	0.4	20	1	AAA55807	Human histone deac	353	15.8	0.4	21	1	ADF75334	Human RT-PCR prime
c 281	16	0.4	0.4	20	1	AAA94502	Antisense oligonuc	354	15.8	0.4	21	1	ADN02584	Primer #2 of the i
c 282	16	0.4	0.4	20	1	AAA94504	Antisense oligonuc	355	15.8	0.4	21	1	ADK61698	Base containing SS
c 283	16	0.4	0.4	20	1	AAA94503	Antisense oligonuc	c 356	15.8	0.4	21	1	ADO11900	Single multiplex P
c 284	16	0.4	0.4	20	1	AAA94505	Antisense oligonuc	c 357	15.6	0.4	30	1	AZ444310	Human SCA7 primer
c 285	16	0.4	0.4	20	1	AAA94506	Antisense oligonuc	c 358	15.6	0.4	30	1	AA513781	Simple sequence re
c 286	16	0.4	0.4	20	1	AAH43117	Antisense oligo, t	c 359	15.4	0.4	17	1	AAT81046	Human c-myb hamme
c 287	16	0.4	0.4	20	1	AAH89537	Human HDAC-2 PCR p	c 360	15.4	0.4	17	1	AAT81049	Human c-myb hamme
c 288	16	0.4	0.4	20	1	AAH89546	Human HDAC-2 antis	c 361	15.4	0.4	17	1	AAT81045	Human c-myb hamme
c 289	16	0.4	0.4	20	1	ABZ85596	Human oligonucleot	c 362	15.4	0.4	17	1	AAT74181	Salmonella enterit
290	16	0.4	0.4	20	1	ABZ88039	Human oligonucleot	c 363	15.4	0.4	17	1	AA336659	PCR primer for mar
291	16	0.4	0.4	20	1	ABD24269	Human calmodulin 2	c 364	15.4	0.4	17	1	AAF01716	Hammerhead ribozym
c 292	16	0.4	0.4	20	1	ABD21826	Human stanniocalci	365	15.4	0.4	17	1	ABK00766	Human NOGO Inozyme
293	15.8	0.4	0.4	19	1	AAZ72847	Human biallelic ma	366	15.4	0.4	17	1	ABK02370	Human NOGO Amberzy
294	15.8	0.4	0.4	19	1	ADL79842	Human HER1 (EGFR)	367	15.4	0.4	17	1	ABK01554	Human NOGO G-Cleav
c 295	15.8	0.4	0.4	19	1	ADL79535	Human HER1 (EGFR)	c 368	15.4	0.4	17	1	ABK00767	Human NOGO Inozyme
c 296	15.8	0.4	0.4	19	1	ADG64260	Y copy of Adican	369	15.4	0.4	17	1	ABK01792	Human NOGO Zinzyme
c 297	15.8	0.4	0.4	19	1	ADH70599	Human Vbeta gene r	c 370	15.4	0.4	17	1	ABK01549	Human NOGO G-Cleav
c 298	15.8	0.4	0.4	20	1	AAQ51743	Mycobacteria probe	371	15.4	0.4	17	1	ABL46975	Human GRID zinzyme
c 299	15.8	0.4	0.4	20	1	AAH8505	S-adenosylmethioni	372	15.4	0.4	17	1	ABN07810	Human GMLP-1 17-m
c 300	15.8	0.4	0.4	20	1	AAV35212	Hepatitis C virus	c 373	15.4	0.4	17	1	ABK19261	Human ERG Amberzym
301	15.8	0.4	0.4	20	1	AAZ33971	Human HG38 DNA PCR	c 374	15.4	0.4	17	1	ABV89508	Human POSHL1 scann
302	15.8	0.4	0.4	20	1	AAZ76854	PCR primer for clo	c 375	15.4	0.4	17	1	ADC37834	Human AMLP1a scann
c 303	15.8	0.4	0.4	20	1	AAZ05303	PCR primer used to	376	15.4	0.4	17	1	ADI51314	Human tumour suppr
c 304	15.8	0.4	0.4	20	1	AAZ93524	PCR primer used to	377	15.4	0.4	17	1	ADM54298	Human GRID mRNA su
c 305	15.8	0.4	0.4	20	1	AAZ97150	PCR primer used to	378	15.4	0.4	17	1	ACN70900	Human GMLP-1 prob
c 306	15.8	0.4	0.4	20	1	AAZ73254	Reverse primer #46	c 379	15.4	0.4	18	1	AAT93486	DOAL allele determ
c 307	15.8	0.4	0.4	20	1	AAZ73052	Human daxe inhibit	380	15.4	0.4	18	1	AAT93485	DOAL allele determ
308	15.8	0.4	0.4	20	1	AAZ24100	Lactococcus lactis	c 381	15.4	0.4	18	1	AAZ90265	DOAL gene PCR prim
c 309	15.8	0.4	0.4	20	1	AAH56611	Streptococcus pyrog	382	15.4	0.4	18	1	AAZ90264	DOAL gene PCR prim
c 310	15.8	0.4	0.4	20	1	AAH89128	Canine retroviral	383	15.4	0.4	18	1	AAH19623	Oligonucleotide co
c 311	15.8	0.4	0.4	20	1	ABK91138	ALS-2 control PCR	c 384	15.4	0.4	18	1	AAH19624	Complementary olig
c 312	15.8	0.4	0.4	20	1	ABQ78562	Primer Rb2 used to	c 385	15.4	0.4	18	1	AAH76247	Human macrophage 1
c 313	15.8	0.4	0.4	20	1	ABQ78586	Primer Rb1 used to	386	15.4	0.4	18	1	ABZ97881	Human UDP-glucuron
314	15.8	0.4	0.4	20	1	ABV99473	Human NOV16a forwa	387	15.4	0.4	18	1	ABA93493	GAGA-B receptor 1a
c 315	15.8	0.4	0.4	20	1	ABV99470	Human NOV16a forwa	388	15.4	0.4	18	1	ABK11198	Oligonucleotide #1
c 316	15.8	0.4	0.4	20	1	ABL50604	Mouse Nafi-7 PCR p	c 389	15.4	0.4	18	1	ABK11199	Oligonucleotide #2
c 317	15.8	0.4	0.4	20	1	AAD39531	Human calreticulin	390	15.4	0.4	18	1	AAD36191	Human Smad6 antis
c 318	15.8	0.4	0.4	20	1	AAK44419	Human HPK/GCK-like	391	15.4	0.4	18	1	ABZ81759	Huntington's disea
c 319	15.8	0.4	0.4	20	1	ABL94386	Mouse C/EBP beta p	c 392	15.4	0.4	18	1	ACC46880	Human COPD related
c 320	15.8	0.4	0.4	20	1	ADG34599	Phosphorothioate o	393	15.4	0.4	18	1	ADK67650	Human COPD related
c 321	15.8	0.4	0.4	20	1	ADA44765	Antisense oligonuc	394	15.4	0.4	18	1	ADK52169	Primer #4 of the i
c 322	15.8	0.4	0.4	20	1	ADH93846	Human gene PCR pri	395	15.4	0.4	18	1	ADN97298	Primer of the inve
c 323	15.8	0.4	0.4	20	1	ABZ88038	Human oligonucleot	396	15.4	0.4	19	1	AAZ10259	PCR primer used to
324	15.8	0.4	0.4	20	1	ABZ88202	Human oligonucleot	c 397	15.4	0.4	19	1	AAA49353	Primer for sequenc
325	15.8	0.4	0.4	20	1	ABD24268	Human calmodulin 2	c 398	15.4	0.4	19	1	AAA49370	Primer for selecti

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OM nucleic - nucleic search, using sw model

Run on: May 12, 2005, 11:24:49 ; Search time 19 Seconds
(without alignments)
3.684 Million cell updates/sec

Title: us-10-029-115-1

Perfect score: 3951

Sequence: 1 gccctatgggacaccgc.....tcataactggtgaaagggc 3951

Scoring table: IDENTITY NUC

Gapop 10.0 , Gapext 0.5

Searched: 428 segs, 8859 residues

Total number of hits satisfying chosen parameters: 856

Minimum DB seq length: 8

Maximum DB seq length: 80

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 439 summaries

Database : rngdb.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	40.4	1.0	51	ADC17063	Human single nucle
2	34	0.9	42	AAT78911	Poly-glutamine rep
3	34	0.9	42	AAS13782	Simple sequence re
4	31.2	0.8	36	AAT78910	Poly-glutamine rep
5	30.2	0.8	35	ABZ81770	Huntington's disea
6	30	0.8	39	ABZ81763	Huntington's disea
7	28.4	0.7	30	AZ44310	Human SCA7 primer
8	28.4	0.7	30	AAS13781	Simple sequence re
9	28.4	0.7	30	ADN97224	AGC1 locus. Unide
10	28.4	0.7	31	AAQ98457	Sense probe CAG-30
11	28.4	0.7	31	AAZ24996	Oligonucleotide CA
12	27.8	0.7	33	ABX79926	EST polymorphic DN
13	26.8	0.7	30	ABZ81777	Huntington's disea
14	24.6	0.6	31	ADE34157	Mink3 Lys54Arg mut
15	23.6	0.6	31	AA131038	Human single nucle
16	23.4	0.6	25	ABZ81767	Huntington's disea
17	23.4	0.6	25	ABZ81768	Huntington's disea
18	23.4	0.6	30	ADE34158	Mink3 Lys54Arg mut
19	23	0.6	23	AAZ40517	Human STE20-relate
20	23	0.6	23	AAZ40518	Human STE20-relate
21	23	0.6	23	ABL39570	Human cancer suppr
22	22.4	0.6	24	ADN97225	Primer of the inve
23	22.4	0.6	24	ADN97164	Primer of the inve
24	22.4	0.6	24	ADR68635	DNA G-quadruplex s
25	22.2	0.6	27	ABK11030	Human HPK/GCK-like
26	22.2	0.6	29	AAA03952	Polymorphic fragme
27	22	0.6	25	ADC38187	Human AMLPia scann
28	22	0.6	25	ADC38189	Human AMLPia scann
29	22	0.6	25	ADC38186	Human AMLPia scann
30	22	0.6	25	ADC38188	Human AMLPia scann
31	21.4	0.5	27	ADO43735	PCR primer used to
32	21	0.5	21	AAZ40515	Human STE20-relate
33	21	0.5	25	ADC38190	Human AMLPia scann

34	21	0.5	25	1	ADC38185	Human AMLPia scann
35	20.4	0.5	22	1	ABX88725	Human Pur alpha an
36	20	0.5	20	1	AAZ40516	Human STE20-relate
37	20	0.5	21	1	AAQ14196	Oligonucleotide pr
38	20	0.5	25	1	ADC38191	Human AMLPia scann
39	20	0.5	25	1	ADC38184	Human AMLPia scann
40	19.8	0.5	24	1	ADJ92110	PCR primer 1 relat
41	19.6	0.5	26	1	ABS71093	Human GPCR ligand
42	19.6	0.5	26	1	ADD69029	Angiogenesis inhib
43	19.4	0.5	21	1	AAF99580	Immunostimulatory
44	19.4	0.5	21	1	ABS78296	Angiogenesis inhib
45	19.4	0.5	21	1	ABL38849	Immunostimulatory
46	19.4	0.5	21	1	ABK10202	Double stranded DN
47	19.4	0.5	21	1	ACH03118	Immunostimulatory
48	19.4	0.5	21	1	ADB37082	Immunostimulatory
49	19.4	0.5	24	1	ABK16111	Porcine GPR8-relat
50	19.4	0.5	24	1	ABK94601	G-protein-coupled
51	19.4	0.5	24	1	ABX92931	Screening method r
52	19.4	0.5	24	1	ADC51835	GPR8 PCR primer, S
53	19	0.5	19	1	AAZ40519	Human STE20-relate
54	19	0.5	20	1	ABZ86076	Human oligonucleot
55	19	0.5	20	1	ABD22306	Human stanniocalci
56	19	0.5	21	1	ADL17771	Human NOV-3 DNA am
57	19	0.5	24	1	ABX03797	DNA encoding secre
58	19	0.5	25	1	ADC38183	Human AMLPia scann
59	19	0.5	25	1	ADC38192	Human AMLPia scann
60	18.8	0.5	23	1	AA157112	Human epithelial c
61	18.8	0.5	24	1	ABN83820	Human prostate-spe
62	18.8	0.5	25	1	ADN62597	Digital karyotypin
63	18.6	0.5	42	1	AAS13782	Simple sequence re
64	18.4	0.5	20	1	AAV52748	Angiotensin-conver
65	18.4	0.5	20	1	AA520967	PCR primer Snrpn-U
66	18.4	0.5	20	1	AAZ37201	Human HMK4 antise
67	18.4	0.5	20	1	ABK44415	Human HPK/GCK-like
68	18.4	0.5	20	1	ABZ86068	Human oligonucleot
69	18.4	0.5	20	1	ABD22298	Human stanniocalci
70	18.4	0.5	20	1	ADH58803	Human CDC-like kin
71	18.4	0.5	20	1	ADH58730	Human CDC-like kin
72	18.4	0.5	23	1	ADD69462	5' anchored (ISSR)
73	18.2	0.5	23	1	AAT77693	Wheat microsatelli
74	18.2	0.5	23	1	AAT96645	Human SCA2 gene PC
75	18.2	0.5	24	1	ABK12118	Human hRDRI RT-PCR
76	18.2	0.5	24	1	ADJ92111	PCR primer 2 relat
77	18.2	0.5	24	1	ADN97247	Primer of the inve
78	18	0.5	18	1	AA631144	Antisense oligonuc
79	18	0.5	18	1	ABZ81780	Huntington's disea
80	18	0.5	18	1	ABZ81779	Huntington's disea
81	18	0.5	18	1	ADS16437	Allele A oligo #2,
82	18	0.5	18	1	ADS16436	Allele A oligo #1,
83	18	0.5	20	1	ABZ31489	Candida albicans G
84	18	0.5	20	1	ADD69519	ISSR-related PCR p
85	18	0.5	42	1	AAT78911	Poly-glutamine rep
86	17.8	0.5	21	1	AAZ61533	Primer 6U for a hu
87	17.8	0.5	21	1	ABZ81769	Huntington's disea
88	17.8	0.5	22	1	ABZ40548	Human ZC1 primer #
89	17.8	0.5	22	1	ABX94818	Human cysteine-ric
90	17.6	0.4	23	1	AAZ85350	Spider silk protei
91	17.4	0.4	19	1	AAT39475	Steroidogenesis ac
92	17.4	0.4	20	1	AAZ55806	Human histone deac
93	17.4	0.4	20	1	AAK94988	Human cDNA clone-s
94	17.4	0.4	20	1	AAH43116	Antisense oligo, t
95	17.4	0.4	20	1	AAH57033	Human estradiol re
96	17.4	0.4	20	1	AAAC89545	Human HDAC-2 antis
97	17.4	0.4	20	1	AAAC89536	Human HDAC-2 PCR p
98	17.4	0.4	20	1	ABK30537	Human glioma-associ
99	17.4	0.4	20	1	ABZ30516	Candida albicans G
100	17.4	0.4	20	1	ABK44442	Human HPK/GCK-like
101	17.4	0.4	20	1	ADD21775	Mouse mdm2 antisen
102	17.4	0.4	20	1	ADL32200	Clone specific PCR
103	17.4	0.4	20	1	ADL11408	Human CDC14A DNA a
104	17.4	0.4	20	1	ADO01250	Human CDC14A antis
105	17.4	0.4	20	1	ADP20520	Transcription fact
106	17.4	0.4	21	1	AAA37188	Human PRO1315 forw

C 253	15.4	0.4	19	1	AR293745	ACCESSION: AR293745
C 254	15.4	0.4	20	1	AR086207	ACCESSION: AR086207
C 255	15.4	0.4	20	1	AR108704	ACCESSION: AR108704
C 256	15.4	0.4	20	1	AR176773	ACCESSION: AR176773
C 257	15.4	0.4	20	1	BD143962	ACCESSION: BD143962
C 258	15.4	0.4	20	1	C0868870	ACCESSION: C0868870
C 259	15.4	0.4	20	1	AR224718	ACCESSION: AR224718
C 260	15.4	0.4	20	1	AR261783	ACCESSION: AR261783
C 261	15.4	0.4	20	1	AR489922	ACCESSION: AR489922
C 262	15.4	0.4	20	1	AX662846	ACCESSION: AX662846
C 263	15.2	0.4	20	1	DOG21144P01	ACCESSION: L78629
C 264	15.2	0.4	20	1	AR103769	ACCESSION: AR103769
C 265	15.2	0.4	20	1	AR117718	ACCESSION: AR117718
C 266	15.2	0.4	20	1	AR121002	ACCESSION: AR121002
C 267	15.2	0.4	20	1	AR129483	ACCESSION: AR129483
C 268	15.2	0.4	20	1	AR130120	ACCESSION: AR130120
C 269	15.2	0.4	20	1	AR130133	ACCESSION: AR130133
C 270	15.2	0.4	20	1	AR159548	ACCESSION: AR159548
C 271	15.2	0.4	20	1	AR162425	ACCESSION: AR162425
C 272	15.2	0.4	20	1	AR163981	ACCESSION: AR163981
C 273	15.2	0.4	20	1	BD230672	ACCESSION: BD230672
C 274	15.2	0.4	20	1	BD250356	ACCESSION: BD250356
C 275	15.2	0.4	20	1	BD272623	ACCESSION: BD272623
C 276	15.2	0.4	20	1	CQ784337	ACCESSION: CQ784337
C 277	15.2	0.4	20	1	AR193126	ACCESSION: AR193126
C 278	15.2	0.4	20	1	AR193131	ACCESSION: AR193131
C 279	15.2	0.4	20	1	AR193159	ACCESSION: AR193159
C 280	15.2	0.4	20	1	AR225891	ACCESSION: AR225891
C 281	15.2	0.4	20	1	AR231469	ACCESSION: AR231469
C 282	15.2	0.4	20	1	AR314465	ACCESSION: AR314465
C 283	15.2	0.4	20	1	AR337050	ACCESSION: AR337050
C 284	15.2	0.4	20	1	AR373625	ACCESSION: AR373625
C 285	15.2	0.4	20	1	AR531371	ACCESSION: AR531371
C 286	15.2	0.4	20	1	AX038745	ACCESSION: AX038745
C 287	15.2	0.4	20	1	AX149138	ACCESSION: AX149138
C 288	15.2	0.4	20	1	AX164704	ACCESSION: AX164704
C 289	15.2	0.4	20	1	AX294455	ACCESSION: AX294455
C 290	15.2	0.4	20	1	AX295117	ACCESSION: AX295117
C 291	15.2	0.4	20	1	AX297351	ACCESSION: AX297351
C 292	15.2	0.4	20	1	AX418622	ACCESSION: AX418622
C 293	15.2	0.4	20	1	AX467415	ACCESSION: AX467415
C 294	15.2	0.4	20	1	AX611052	ACCESSION: AX611052
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ACCESSION
184400
VERSION
184400.1 GI:3021920
KEYWORDS
SOURCE
Unknown.
ORGANISM
Unknown.
REFERENCE
1 (bases 1 to 51)
AUTHORS
Schalling,M., Hudson,T.J. and Housman,D.E.
TITLE
Direct detection of expanded nucleotide repeats in the human genome
JOURNAL
Patent: US 5695933-A 1 09-DEC-1997;
FEATURES
Location/Qualifiers
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source

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DB	51	CAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAG	1
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DEFINITION	Sequence 6 from Patent WO9717445.		
ACCESSION	A62705		
VERSION	A62705.1	GI:3716589	
KEYWORDS			
SOURCE	unidentified		
ORGANISM	unidentified		
REFERENCE	unclassified.		
AUTHORS	1		
TITLE	Tora, L., Lutz, Y., Trottier, Y., Mandel and Jean-Louis.		
	METHOD FOR TREATING NEURODEGENERATIVE DISEASES USING A 1C2 ANTIBODY		
	OR A FRAGMENT OR DERIVATIVE THEREOF, AND CORRESPONDING		
	PHARMACEUTICAL COMPOSITIONS		
JOURNAL	Patent: WO 97/17445-A 6 15-MAY-1997;		
COMMENT	CENTRE NAT RECH SCIENT (FR)		
FEATURES	Other publication FR 2741088 19970516.		
	Location/Qualifiers		
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VERSION	A62704.1	GI:3716588			
KEYWORDS	.				
SOURCE	unidentified				
ORGANISM	unidentified				
	unclassified.				

1	Tora, L., Lutz, Y., Trotter, Y., Mandel and Jean-Louis.
REFERENCE	METHOD FOR TREATING NEURODEGENERATIVE DISEASES USING A 1C2 ANTIBODY
AUTHORS	OR A FRAGMENT OR DERIVATIVE THEREOF, AND CORRESPONDING
TITLE	PHARMACEUTICAL COMPOSITIONS
JOURNAL	Patent: WO 9717445-A 5 15-MAY-1997;
COMMENT	CENTRE NAT RECH SCIEN (FR)
FEATURES	Other publication FR 2741088 19970516.
source	Location/Qualifiers
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May

GenCore version 5.1.1.6
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Sequence Search Alignment

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LOCUS AR107612 20 bp DNA linear PAT 14-FEB-2001
DEFINITION Sequence 52 from patent US 6110664.
ACCESSION AR107612
VERSION AR107612.1 GI:12823099
KEYWORDS
SOURCE Unknown.
ORGANISM Unknown.
Unclassified.
REFERENCE 1 (bases 1 to 20)
AUTHORS Cowser, L.M.
TITLE Antisense inhibition of G-alpha-S1 expression
JOURNAL Patent: US 6110664-A 52 29-AUG-2000;
FEATURES Location/Qualifiers
source 1..20
/organism="unknown"
/mol_type="unassigned DNA"

Query Match 0.4%; Score 16; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.7e+02;
Matches 16; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1112 TAAACAGCAGCAGCA 1127
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Db 16 TAAACAGCAGCAGCA 1

Sequence search alignment

RESULT 26
AR453231/c
LOCUS AR453231 23 bp DNA linear PAT 20-FEB-2004
DEFINITION Sequence 60 from patent US 6680170.
ACCESSION AR453231
VERSION AR453231.1 GI:42685485
KEYWORDS
SOURCE Unknown.
ORGANISM Unknown.
Unclassified.
REFERENCE 1 (bases 1 to 23)
AUTHORS Plowman,G., Martinez,R. and Whyte,D.
TITLE Polynucleotides encoding STE20-related protein kinases and methods
of use
JOURNAL Patent: US 6680170-A 60 20-JAN-2004;
FEATURES Location/Qualifiers
source 1..23
/organism="unknown"
/mol_type="genomic DNA"

Query Match 0.6%; Score 23; DB 1; Length 23;
Best Local Similarity 100.0%; Pred. No. 35;
Matches 23; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3454 ACAGTAGAGGAGGGGCAGCGGCT 3476
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Db 23 ACAGTAGAGGAGGGGCAGCGGCT 1